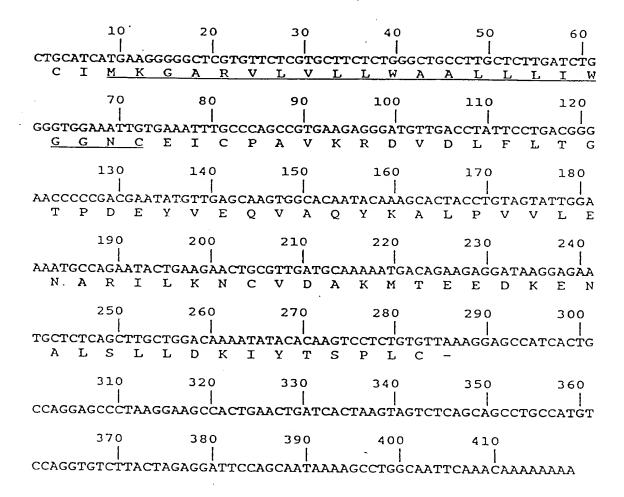
#### F19.1

FIGURE 1.
TRFP CHAIN 1, LEADER A



#### F19. 2

FIGURE 2.
TRFP CHAIN 1, LEADER B

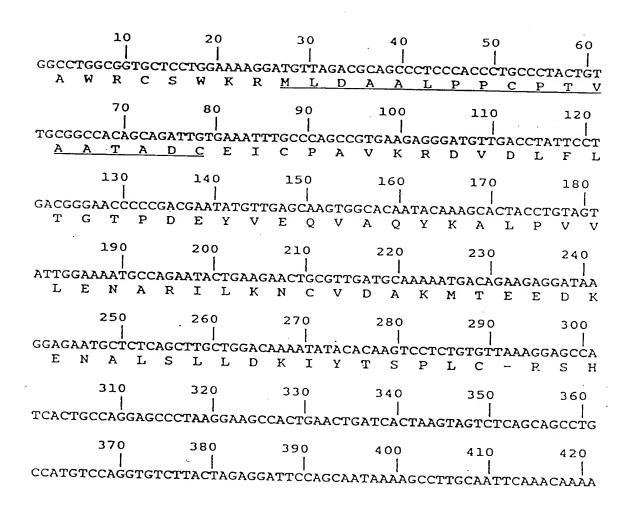


FIGURE 3.
TRFP CHAIN 2, LONG FORM

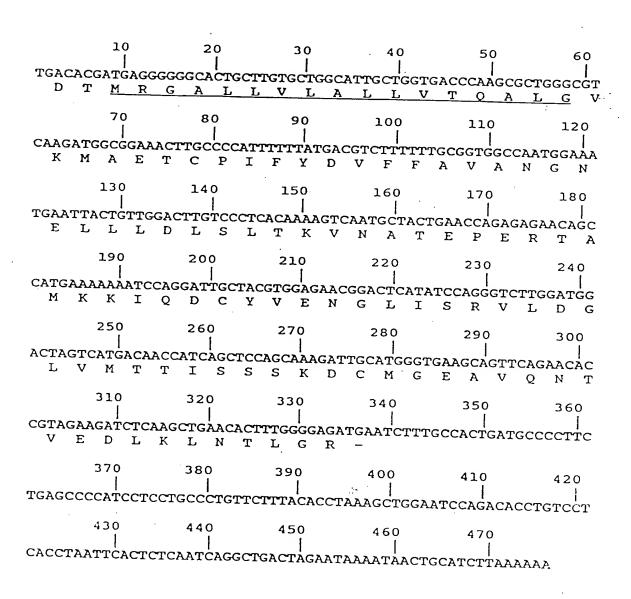


FIGURE 4. TRFP I CHAIN 2, SHORT FORM

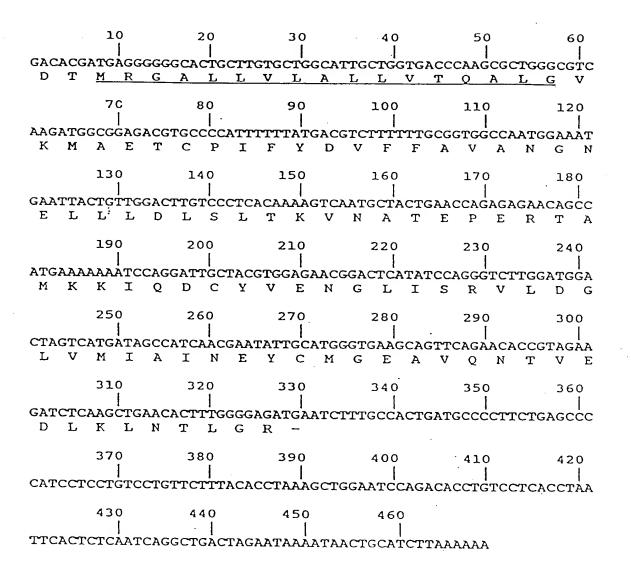
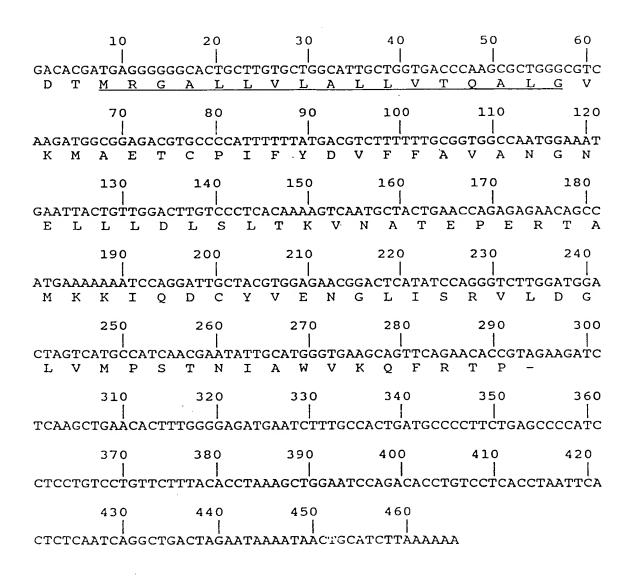


FIGURE 5.
TRFP CHAIN 2, TRUNCATED SHORT FORM



# FIGURE 6. TREP CHAIN #1 PROTEIN SEQUENCE

-20 CIMKGARVLVLLHAALLLIWGGNC	AWRCSWKRMLDAALPPCPTBAATADC
Cl Leader A	Cl Leader B

35	T A A	1
30	YKALP	1 1 1 1 1 1
25	EQVAQ	1
20	PDEYV	1 1 1
15	FLIGI	1 1 1
10	RDVDL	1 1 1 1
 	EICPAVKRDVDLFLTGTPDEYVEQVAQYKALPVV	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
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FIGURE 7.
TREP CHAIN #2 PROTEIN SEQUENCES

-10 DTMRGALLVLALLVTQALG	5 10 15 20 25 30 35 40 VKMAETCPIFYDV <b>FFAVA</b> NGNELLLDLSLTKV <u>NA</u> TEPERT	1	1	1
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# FIG. 7 (con't.)

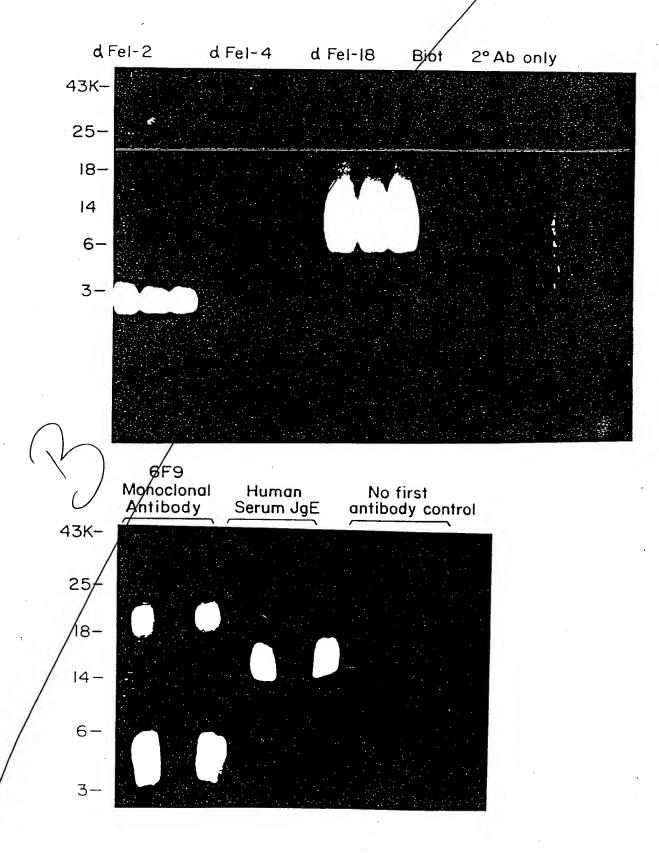
FIGURE 7 (continued)

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85 90 C2L TVEDLKLNTLG

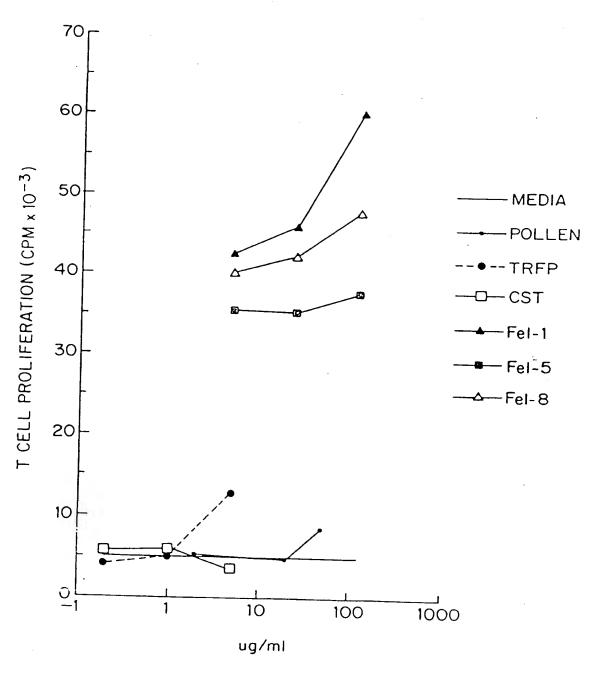
ZRO. T V

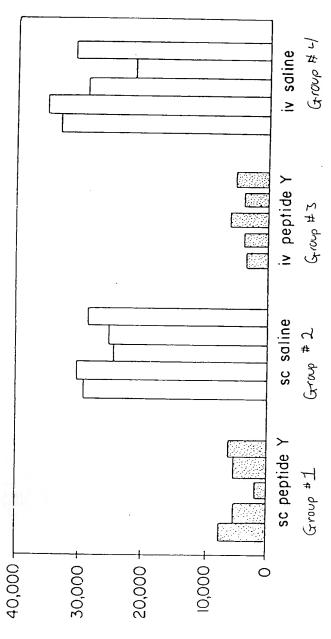
c2s



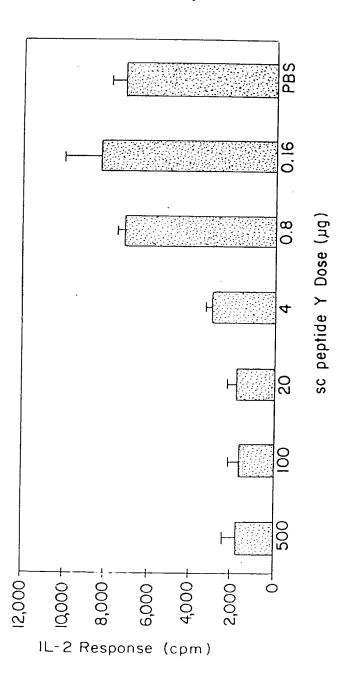


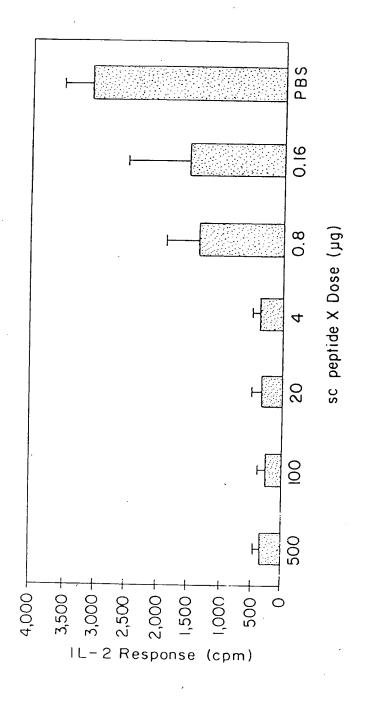
PATIENT #131.2 2° (TRFP:1°)

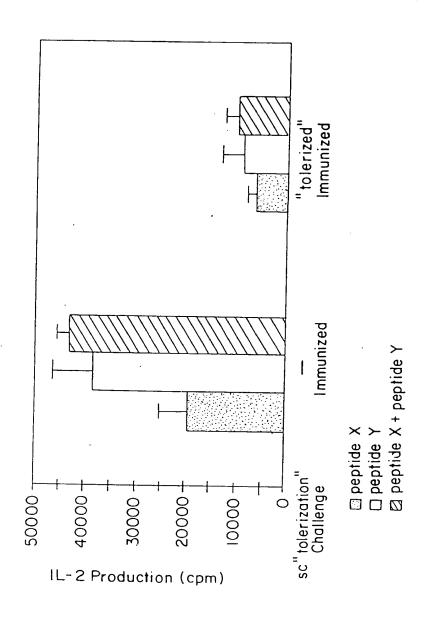


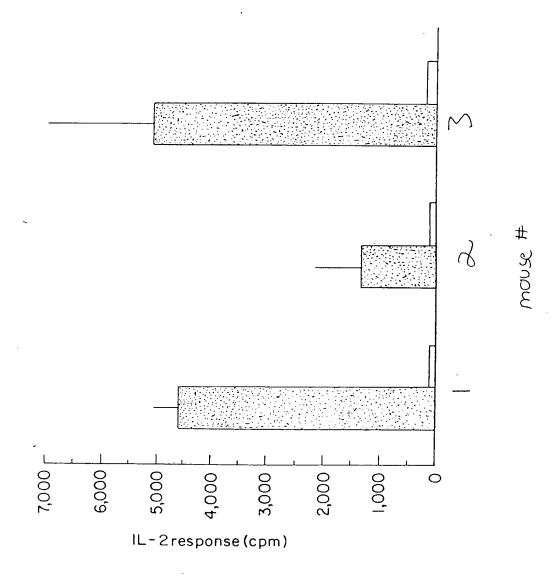


IL-2 Production (cpm)







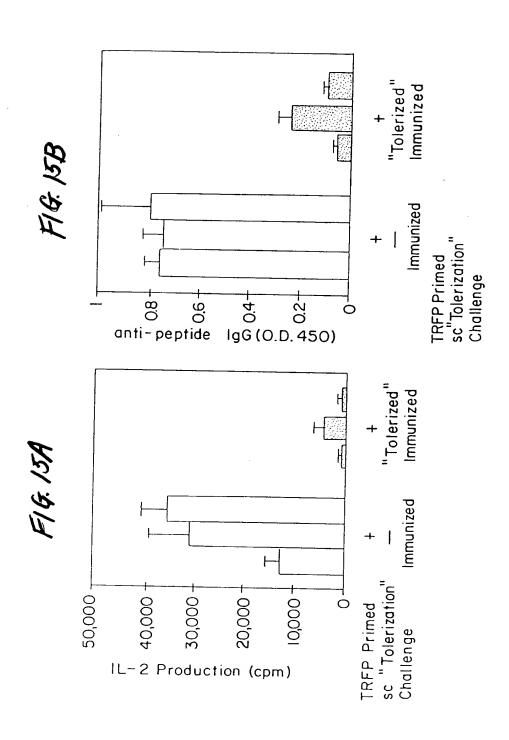


[ - (ultured without Poptide

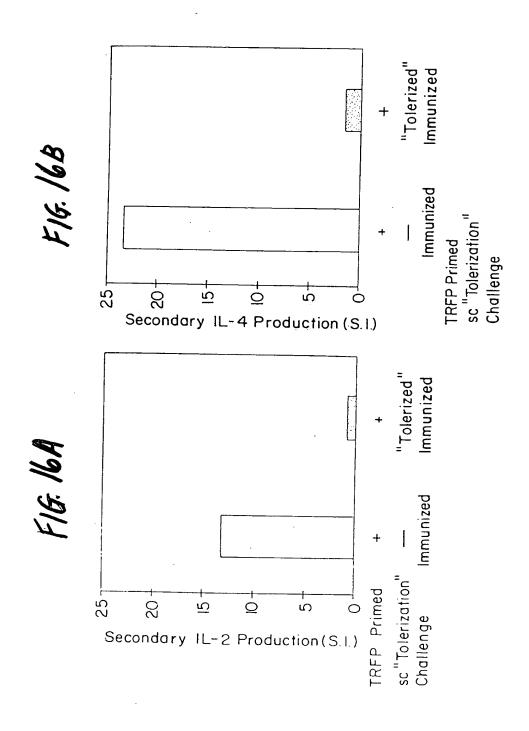
1 - Cultured with Peptide

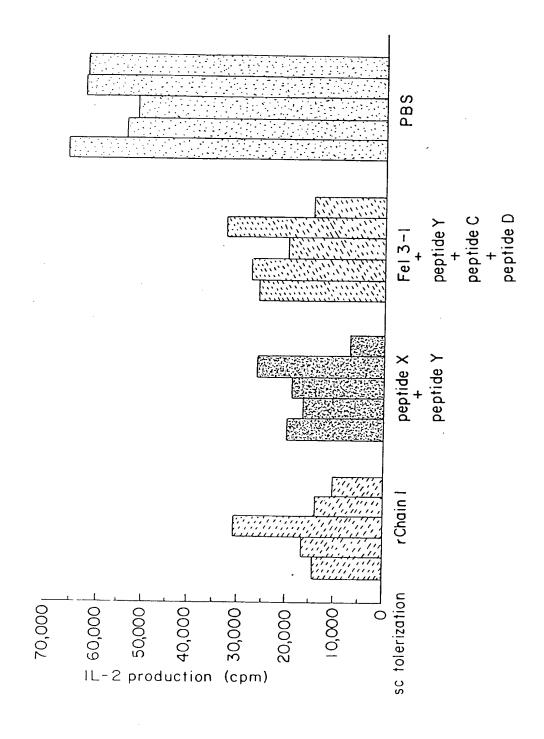
TEGEND

FIG. 15A \$ 15B



#### FIG. 16 A \$ 16B





#### Sequence

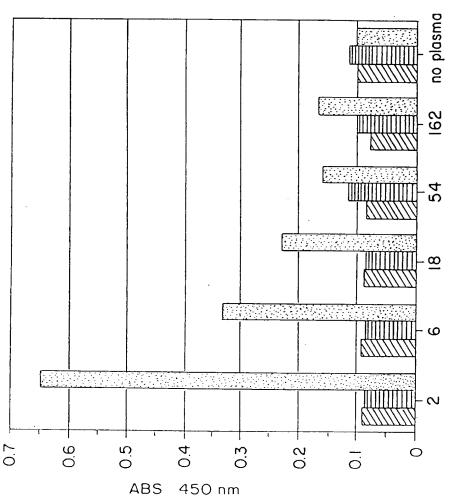
peptide	
name	

X	KRDVDLFLTGTPDEYVEQVAQYKALPV
---	-----------------------------

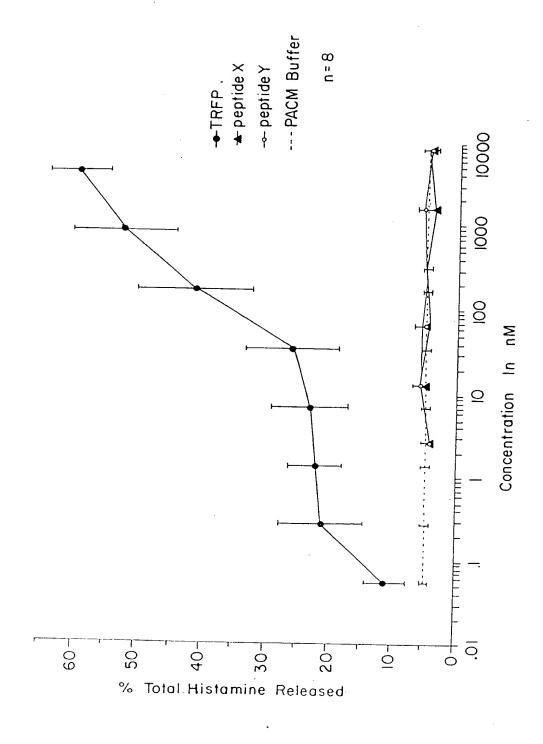
- Y KALPVVLENARILKNCVDAKMTEEDKE
- Z FFAVANGNELLLDLSLTKVNATEPER
- A EEDKENALSLLDKIYTSPL
- B MGEAVQNTVEDLKLNTLGR
- C EEDKENALSLLDKIYT
- D NALSLLDKIYTSPL

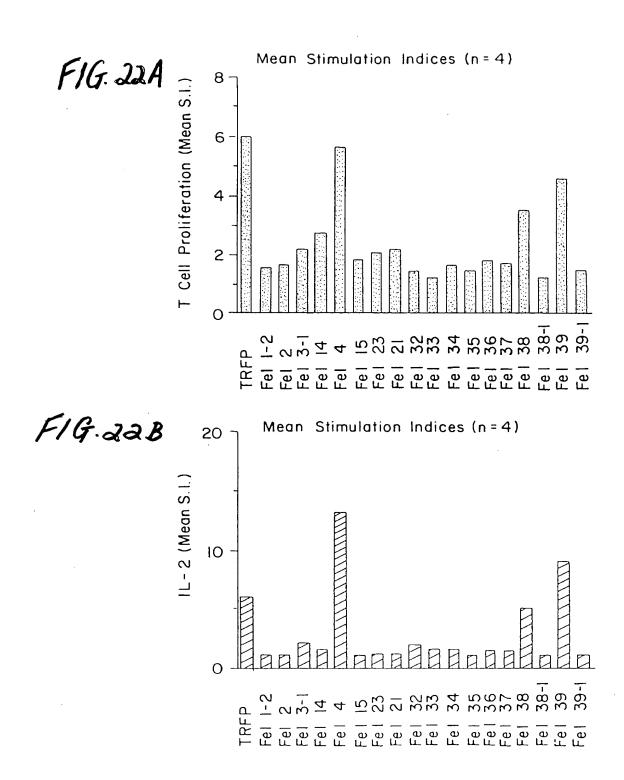
- Fel 32 VKMAETCPIFYDVFFAVA
- FYDVFFAVANGNELLLD Fel 33
- Fel 34 NGNELLLDLSLTKVNATE
- Fel 35 SLTKVNATEPERTAMKKI
- Fel 36 ERTAMKKIQDCYVENGL
- Fel 37 QDCYVENGLISRVLDGLV
- Fel 38 ISRVLDGLVMTTISSSKDCM
- Fel 38-I ISRVLDGLVMIAINE \*\* DCM
- Fel 39 MTTISSSKDCMGEAVQNTEVELDKLNTLGF Fel 39.I MIAINE \*\* DCMGEAVQNTEVELDKLNTLGF

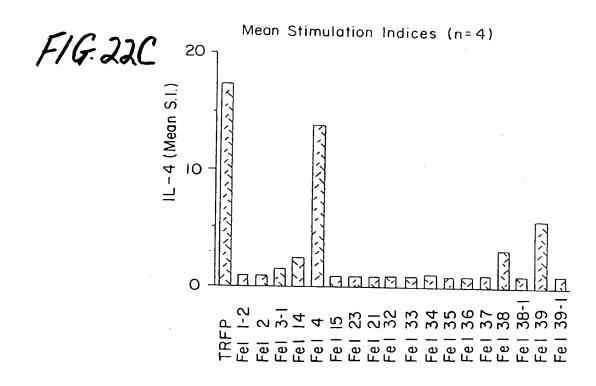
Ist antibody: Pooled Human Plasma (IgG depleted)

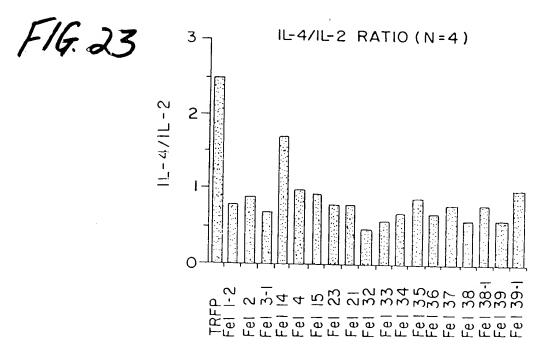


1/dilution

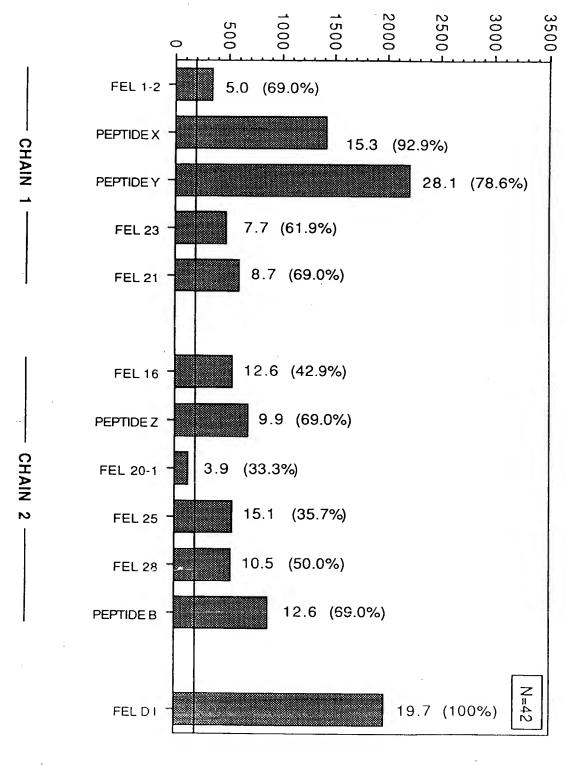


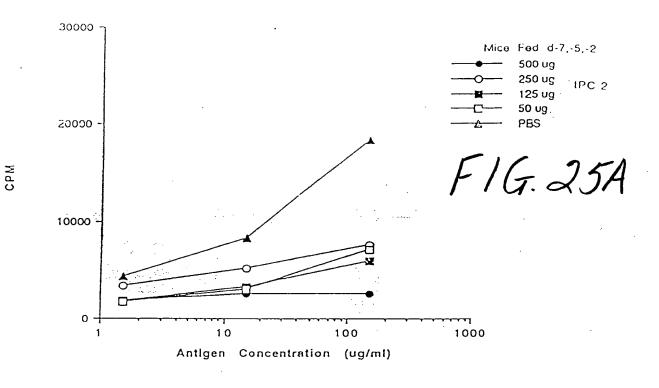


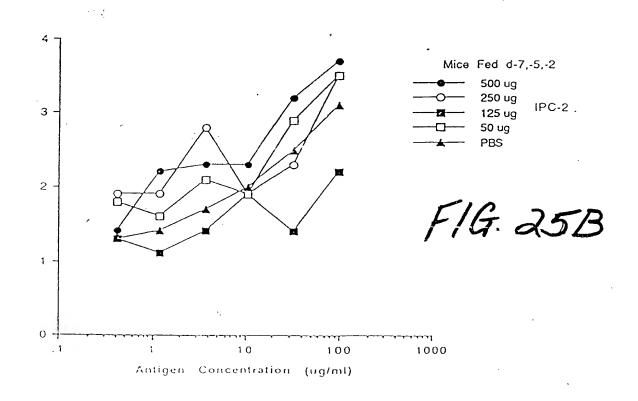




#### POSITIVITY INDEX



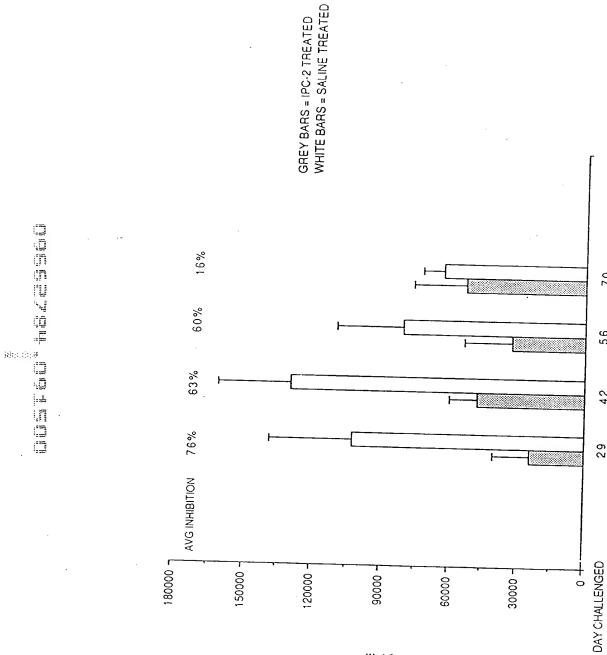




70

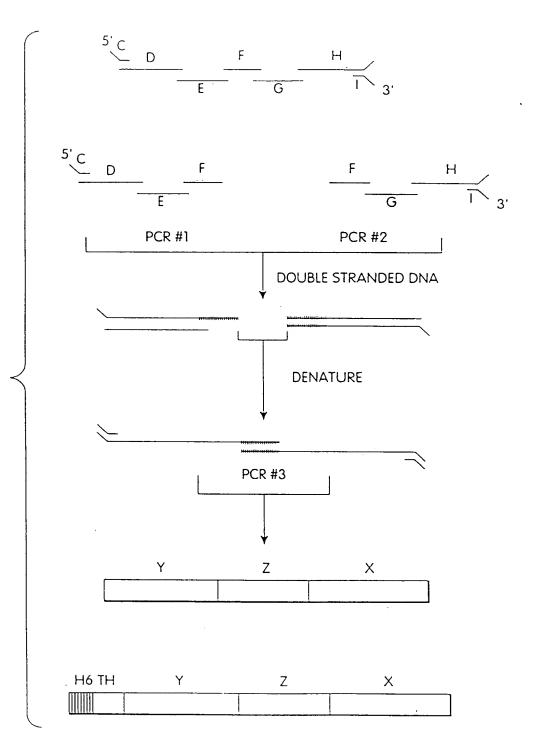
56

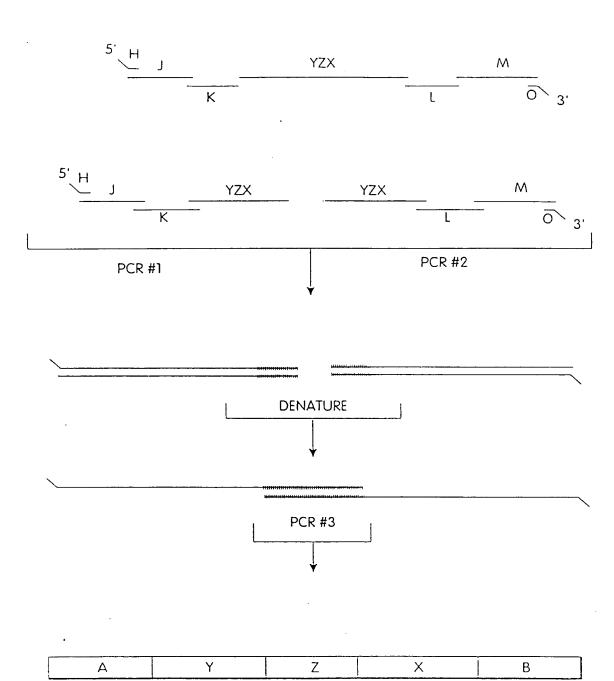
42



СРМ

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BAM HI GGGGGATCCAAAGCTCTGCCGGTTGTT 3' K A' L P V V
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GAAGACAAAGAA 3'

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-ო CAGAACACCGTTGAAGACCTGAAACTGAACACCCTGGGTCGTTGAATGTAACTGCAGAATTCCCC PST I ECORI GACAGAGACGACCTGTTTTAGATGTGGAGAGGCGACTTTCGAGACGGCCAACAAGACCTT CGAGTCATGTTTCGAGACGGCCAATACCCACTTCGACAAGTCTTGTGGCAACTT 5' O T N Q 5 SPLKAL CGAGACGGCCAAATCATCAGATCTGACGTCTTCGAACCTAGGGG - XBAI PSTI HINDIII ECORI GGGGATCCGAAGAAGAAAACGCTCTGTCTCTGCTG 3' AV ريا -TGAAACCCCTCTACTTACATTGACGTCTTAAGGGG 回 G N A D L K L N K A L P V M 回 GGGGATCCGAAGAGACAAA 3' × 口 回 Ω 囶 臼 臼 H G BAM HI. BAM HI <u>-</u> 5 ريا -H Ξ  $\equiv$ 0 ×

ATGGGTCACCACCACCACGAATTCCTGGTTCCGCGTGGATCC

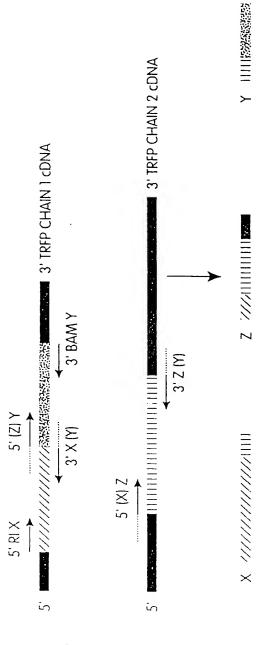
AAAGCTCTGCCGGTTGTTCTGGAAAACGCTCGTATCCTGAAAAACTGC

GTTGACGCTAAAATGACCGAAGAAGAAAGAATTCTTCGCTGTTGCT V D A K M T E E D K E F F A V A AACGGTAACGAACTGCTGGACCTGTCTGACCAAAGTTAACGCT N G N E L L D L S L T K V N A ACCGAACCGGAACGTAACGTTGACCTGTTCCTGACCGGTACC

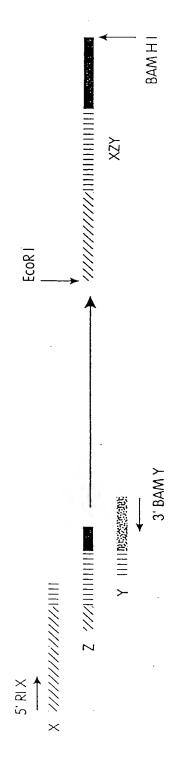
CCGGACGAATACGTTGAACAGGTTGCTCAGTACAAAGCTCTGCCGGTT PDEYVEQVAAQYKALPV

# F/G. 31

# 1) PCR INDIVIDUAL EPITOPES



# 2) LINK ISOLATED EPITOPES BY POOLING/PCR



#### 5' PRIMERS

#### XZY CONSTRUCT

- K R D V D L
  5' XRI 5'-GGGGAATTCAAGAGGGATGTTGACCTA-3'
  ECOR I X
- L P V F F A V A N
  5'(X) Z 5'-CTACCTGTATTTTTTGCGGTGGCCAAT-3'
  X Z
- PERKALPVV
  5'(Z)Y5'-CCAGAGAGAAAGCACTACCTGTAGTA-3'
  Z Y

#### YXZ CONSTRUCT

- K A L P V V
  5' YRI 5'-GGGGAATTCAAAGCACTACCTGTAGTA-3'
  ECOR I Y
- D K E K R D V D L
  5' (Y) X 5'-GATAAGGAGAAGAGGGATGTTGACCTA-3'
  Y X
- L P V F F A V A N
  5' (X) Z 5'-CTACCTGTATTTTTGCGGTGGCCAAT-3'
  X Z

#### ZXY CONSTRUCT

- F F A V A N G
  5' ZRI 5'-GGGGAATTCTTTGCGGTGGCCAATGGA-3'
  ECOR I Z
- K R D V D L P
  5' (Z) X 5'-AAGAGGGATGTTGACCTATTC-3'
  X

# F/G. 32 (cont.)

#### 3' PRIMERS

#### XZY CONSTRUCT

- αΝ αΑ αν αΑ αF αF αν αP αL αΑ αΚ αΥ

  3' Χ (Ζ) 5'-ATTGGCCACCGCAAAAAATACAGGTAGTGCTTTGTA-3'

  Z | Χ

  αL αΑ αΚ αR αΕ αΡ αΕ αΤ αΑ

  3' Ζ (Υ) 5'-TAGTGCTTTTCTCTCTGGTTCAGTAGC-3'

  Υ | Ζ
- αSTOPαE αΚ αD αΕ αΕ αΤ 3' Y BAM 5'-GGGGATCCTTACTCCTTATCCTCTTCTGT-3' BAMH I Y

#### YXZ CONSTRUCT

- αL αD αV αD αR αK αΕ αΚ αD αΕ αΕ αΤ

  3' Y (X) 5'-TAGGTCAACATCCCTCTTCTCCTTATCCTCTTCTGT-3'

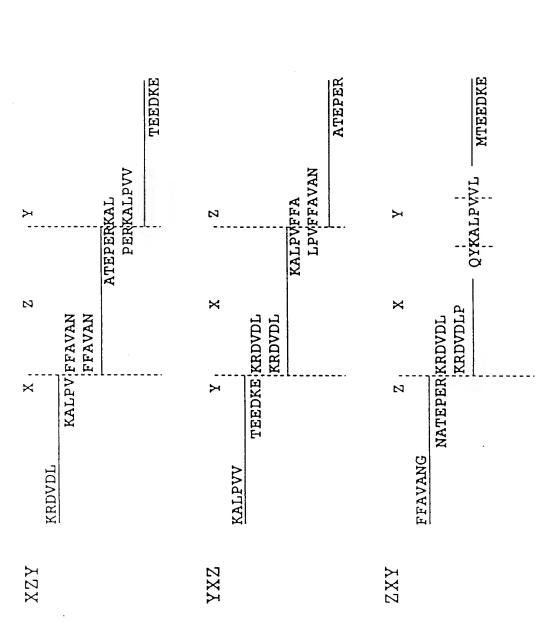
  X Y

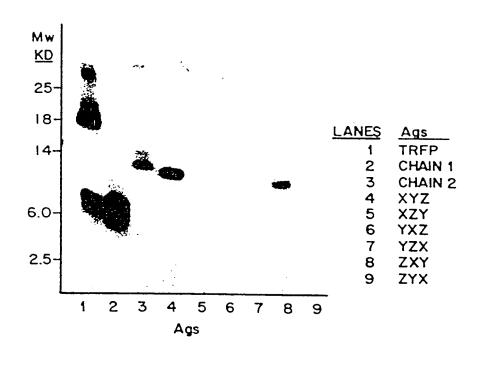
  αA αF αF αV αΡ αL αA αΚ αΥ

  3' X (Z) 5'-CGCAAAAAATACAGGTAGTGCTTTGTA-3'

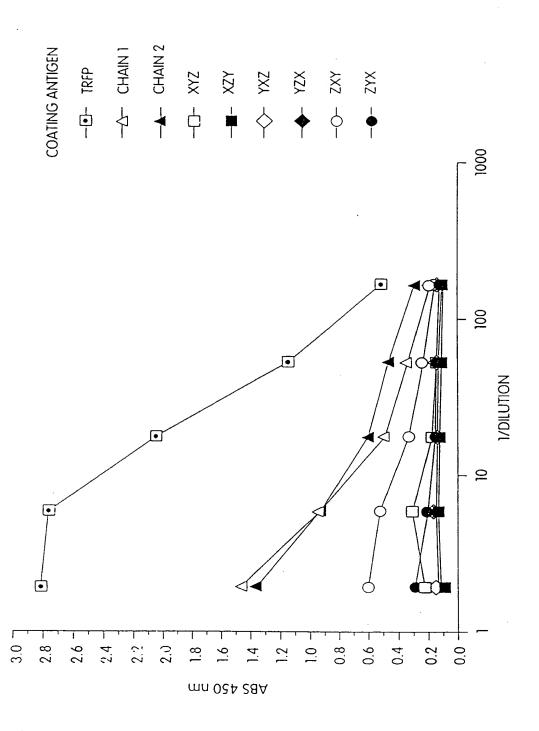
  Z X

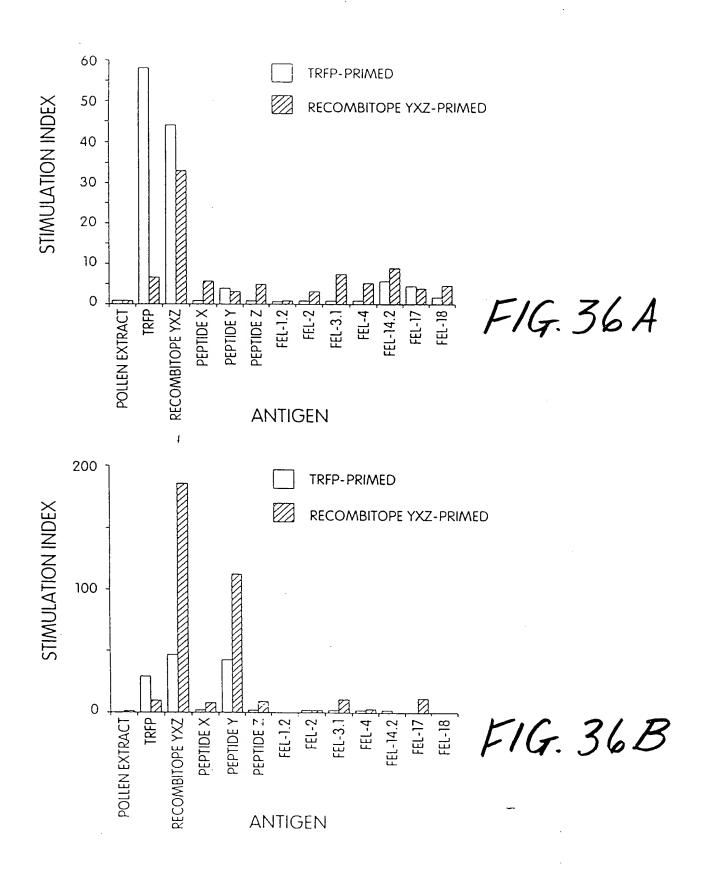
#### **7XY CONSTRUCT**





# F/G. 35





# F1G. 36C

